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IN THE CLAIMS

1. (Original) A composition for inhibiting growth of chondrosarcoma cells comprising an effective amount of a peptide of formula I and a pharmaceutically acceptable carrier:

$$Xaa_{1}-Xaa_{2}-Xaa_{3}-Xaa_{4}-Xaa_{5}-Xaa_{6} Xaa_{7}-Xaa_{8}-Xaa_{9}$$
 (I)

wherein:

Xaa₁, Xaa₄, and Xaa₆ are separately each apolar amino acids;

Xaa2 is a basic amino acid;

Xaa3 is a cysteine-like amino acid;

Xaa₅ is a polar or aliphatic amino acid;

Xaa7 is an acidic amino acid;

Xaa₈ is an aliphatic or polar amino acid; and

Xaa₉ is an aliphatic, apolar or basic amino acid.

2. (Original) A composition for inhibiting growth of chondrosarcoma cells comprising an effective amount of a peptide of formula II and a pharmaceutically acceptable carrier:

$$Xaa_{10}-Xaa_{11}-Xaa_{12} Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (II)

wherein:

Xaa₁₀ is a polar, acidic, basic or apolar amino acid;

Xaa11 is a polar or aromatic amino acid;

Xaa₁₂ is a polar, basic, aliphatic or apolar amino acid;

Xaa₁₃ is an aromatic, aliphatic, polar or acidic amino acid;

Xaa₁₄ is an aromatic, apolar or polar amino acid;

Xaa₁₅ is an apolar or acidic amino acid;

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Xaa₁₆ is a basic, a polar or an apolar amino acid;

Xaa₁₇ is a basic, a polar, an aliphatic, an apolar or an acidic amino acid;

Xaa₁₈ is an apolar or an aliphatic amino acid; and

Xaa₁₉ is a basic or an aliphatic amino acid.

3. (Original) A composition for inhibiting growth of chondrosarcoma cells comprising an effective amount of a peptide of formula III and a pharmaceutically acceptable carrier:

$$Xaa_{1}-Xaa_{2}-Xaa_{3}-Xaa_{4}-Xaa_{5}-Xaa_{6}-Xaa_{7}-Xaa_{8}-Xaa_{9}-Xaa_{10}-Xaa_{11}-Xaa_{12}-Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (III)

wherein

Xaa₁, Xaa₄, and Xaa₆ are separately each apolar amino acids;

Xaa2 is a basic amino acid;

Xaa₃ is a cysteine-like amino acid;

Xaa₅ is a polar or aliphatic amino acid;

Xaa₇ is an acidic amino acid;

Xaa₈ is an aliphatic or polar amino acid;

Xaao is an aliphatic, apolar or basic amino acid;

Xaa₁₀ is a polar, acidic, basic or apolar amino acid;

Xaa11 is a polar or aromatic amino acid;

Xaa₁₂ is a polar, basic, aliphatic or apolar amino acid;

Xaa₁₃ is an aromatic, aliphatic, polar or acidic amino acid;

Xaa₁₄ is an aromatic, apolar or polar amino acid;

Xaa₁₅ is an apolar or acidic amino acid;

Xaa₁₆ is a basic, a polar or an apolar amino acid;

Xaa₁₇ is a basic, a polar, an aliphatic, an apolar or an acidic amino acid;

Xaa₁₈ is an apolar or an aliphatic amino acid; and

Xaa₁₉ is a basic or an aliphatic amino acid.

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4. (Original) A composition for inhibiting growth of chondrosarcoma cells comprising an effective amount of a peptide of formula IV (SEQ ID NO:18) and a pharmaceutically acceptable carrier:

$$Xaa_a-Xaa_b-Xaa_c-Xaa_d-Xaa_e-Xaa_f-Xaa_g-Xaa_h-Xaa_i-Xaa_j-Xaa_k-Xaa_L-Xaa_m-Xaa_n-Xaa_0-Xaa_p-Xaa_1-Xaa_2-Xaa_3-Xaa_4-Xaa_5-Xaa_6-Xaa_7-Xaa_8-Xaa_9-Xaa_{10}-Xaa_{11}-Xaa_{12}-Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (IV)

Xaa_a is proline;

Xaa_b is glutamine or glutamic acid;

Xaac is threonine;

Xaa_d is glycine;

Xaae is aspartic acid or glutamic acid;

Xaa_f is leucine;

Xaag is aspartic acid;

Xaa_h is glutamine or serine;

Xaa_i is asparagine or alanine;

Xaa_i is threonine;

Xaa_k is isoleucine or leucine;

Xaa_l is glutamic acid or lysine;

Xaa_m is threonine or alanine;

Xaa_n is methionine;

Xaa_o is arginine;

Xaa_p is lysine or threonine;

Xaa₁₇ is lysine or aspartic acid;

Xaa₁₉ is lysine.

Xaa₁ is proline;

Xaa₂ is arginine;

Xaa₃ is cysteine;

Xaa₄ is glycine;

Xaa₅ is valine or asparagine;

Xaa₆ is proline;

Xaa₇ is aspartic acid;

Xaa₈ is valine or leucine;

Xaa₉ is alanine or glycine;

Xaa₁₀ is asparagine or arginine;

Xaa₁₁ is tyrosine or phenylalanine;

Xaa₁₂ is asparagine or glutamine;

Xaa₁₃ is phenylalanine or threonine;

Xaa₁₄ is phenylalanine;

Xaa₁₅ is proline or glutamic acid;

Xaa₁₆ is arginine or glycine;

Xaa₁₈ is proline or leucine; and

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5. (Original) The composition of any one of claims 1-4, wherein an apolar amino acid is methionine, glycine or proline.

- 6. (Original) The composition of any one of claims 1-4, wherein a basic amino acid is histidine, lysine, arginine, 2,3-diaminopropionic acid, ornithine, homoarginine, ñ-aminophenylalanine, and 2,4-diaminobutyric acid. The composition of any one of claims 1-4, wherein a cysteine-like amino acid is cysteine, homocysteine, penicillamine, or â-methyl cysteine.
- 7. (Original) The composition of any one of claims 1-4, wherein an aliphatic amino acid is alanine, valine, leucine, isoleucine, t-butylalanine, t-butylalanine, N-methylisoleucine, norleucine, N-methylvaline, cyclohexylalanine, â-alanine, N-methylglycine, or á-aminoisobutyric acid.
- 8. (Original) The composition of any one of claims 1-4, wherein an acidic amino acid is aspartic acid or glutamic acid.
- 9. (Original) The composition of any one of claims 1-4, wherein a polar amino acid is asparagine, glutamine, serine, threonine, tyrosine, citrulline, N-acetyl lysine, methionine sulfoxide, or homoserine, or an apolar amino acid such as methionine, glycine or proline.
- 10. (Original) The composition of any one of claims 1-4, wherein an aromatic amino acid is phenylalanine, tyrosine, tryptophan, phenylglycine, naphthylalanine, â-2-thienylalanine, 1,2,3,4-tetrahydro-isoquinoline-3-carboxylic acid, 4-chlorophenylalanine, 2-fluorophenylalanine, 3-fluorophenylalanine, 4-fluorophenylalanine, pyridylalanine, or 3-benzothienyl alanine.

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11. (Original) The composition of any one of claims 1-4 wherein the peptide inhibits proteinase activity of matrix metalloproteinase-1, matrix metalloproteinase-2, matrix metalloproteinase-3, matrix metalloproteinase-4, matrix metalloproteinase-5, matrix metalloproteinase-6, matrix metalloproteinase-7, matrix metalloproteinase-8, and matrix metalloproteinase-9, matrix metalloproteinase-10, matrix metalloproteinase-11, matrix metalloproteinase-12, or matrix metalloproteinase-13.

- 12. (Original) The composition of any one of claims 1-4 wherein inhibiting growth of chondrosarcoma inhibits growth of conventional chondrosarcoma, myxoid chondrosarcoma, mesenchymal chondrosarcoma, clear cell chondrosarcoma, or dedifferentiated (spindle cell) chondrosarcoma.
- 13. (Original) The composition of any one of claims 1-4 wherein inhibiting growth of chondrosarcoma cells inhibits growth of a bone tumor.
- 14. (Original) The composition of any one of claims 1-4, wherein inhibiting growth of chondrosarcoma cells diminishes a size of a bone tumor.
- 15. (Original) The composition of claim 12, 13 or 14, wherein the tumor is metastatic, non-metastatic, vascularized, non-vascularized, hard or soft.
- (Original) The composition of any one of claims 1-4 wherein the peptide comprises SEQ ID NO:1, SEQ ID NO:2, SEQ ID NO:3, SEQ ID NO:4, SEQ ID NO:5, SEQ ID NO:6, SEQ ID NO:7, SEQ ID NO:8, SEQ ID NO:9, SEQ ID NO:10, SEQ ID NO:11, SEQ ID NO:12, or SEQ ID NO:13.
- 17. (Original) An anti-sarcoma composition that comprises a therapeutically effective amount of peptide that comprises SEQ ID NO:1, SEQ ID NO:2, SEQ ID

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NO:3, SEQ ID NO:4, SEQ ID NO:5, SEQ ID NO:6, SEQ ID NO:7, SEQ ID NO:8, SEQ ID NO:9, SEQ ID NO:10, SEQ ID NO:11, SEQ ID NO:12, or SEQ ID NO:13, and a pharmaceutically acceptable carrier, wherein the peptide is capable of inhibiting growth of chondrosarcoma cells.

18. (Original) A method for decreasing growth of chondrosarcoma cells that comprises contacting a chondrosarcoma cell with an effective amount of a peptide of formula I:

$$Xaa_1-Xaa_2-Xaa_3-Xaa_4-Xaa_5-Xaa_6 Xaa_7-Xaa_8-Xaa_9$$
 (I)

wherein:

Xaa₁, Xaa₄, and Xaa₆ are separately each apolar amino acids;

Xaa₂ is a basic amino acid;

Xaa₃ is a cysteine-like amino acid;

Xaa₅ is a polar or aliphatic amino acid;

Xaa₇ is an acidic amino acid;

Xaa₈ is an aliphatic or polar amino acid; and

Xaa₉ is an aliphatic, apolar or basic amino acid.

19. (Original) A method for decreasing growth of chondrosarcoma cells that comprises contacting a chondrosarcoma cell with an effective amount of a peptide of formula II:

$$Xaa_{10}-Xaa_{11}-Xaa_{12} Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (II)

wherein:

Xaa₁₀ is a polar, acidic, basic or apolar amino acid;

Xaa11 is a polar or aromatic amino acid;

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Xaa₁₂ is a polar, basic, aliphatic or apolar amino acid;

Xaa₁₃ is an aromatic, aliphatic, polar or acidic amino acid;

Xaa₁₄ is an aromatic, apolar or polar amino acid;

Xaa₁₅ is an apolar or acidic amino acid;

Xaa₁₆ is a basic, a polar or an apolar amino acid;

Xaa₁₇ is a basic, a polar, an aliphatic, an apolar or an acidic amino acid;

Xaa₁₈ is an apolar or an aliphatic amino acid; and

Xaa₁₉ is a basic or an aliphatic amino acid.

20. (Original) A method for decreasing growth of chondrosarcoma cells that comprises contacting a chondrosarcoma cell with an effective amount of a peptide of formula III:

$$Xaa_{1}-Xaa_{2}-Xaa_{3}-Xaa_{4}-Xaa_{5}-Xaa_{6}-Xaa_{7}-Xaa_{8}-Xaa_{9}-Xaa_{10}-Xaa_{11}-Xaa_{12}-Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (III) wherein:

Xaa₁, Xaa₄, and Xaa₆ are separately each apolar amino acids;

Xaa2 is a basic amino acid;

Xaa3 is a cysteine-like amino acid;

Xaa₅ is a polar or aliphatic amino acid;

Xaa₇ is an acidic amino acid;

Xaa₈ is an aliphatic or polar amino acid;

Xaa₉ is an aliphatic, apolar or basic amino acid;

Xaa₁₀ is a polar, acidic, basic or apolar amino acid;

Xaa₁₁ is a polar or aromatic amino acid;

Xaa₁₂ is a polar, basic, aliphatic or apolar amino acid;

Xaa₁₃ is an aromatic, aliphatic, polar or acidic amino acid;

Xaa₁₄ is an aromatic, apolar or polar amino acid;

Xaa₁₅ is an apolar or acidic amino acid;

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Xaa₁₆ is a basic, a polar or an apolar amino acid;

Xaa₁₇ is a basic, a polar, an aliphatic, an apolar or an acidic amino acid;

Xaa₁₈ is an apolar or an aliphatic amino acid; and

Xaa₁₉ is a basic or an aliphatic amino acid.

21. (Original) A method for decreasing growth of chondrosarcoma cells that comprises contacting a chondrosarcoma cell with an effective amount of a peptide of formula IV (SEQ ID NO:18):

$$Xaa_a-Xaa_b-Xaa_c-Xaa_d-Xaa_e-Xaa_f-Xaa_g-Xaa_h-Xaa_i-Xaa_j-Xaa_k-Xaa_L-Xaa_m-Xaa_n-Xaa_0-Xaa_p-Xaa_1-Xaa_2-Xaa_3-Xaa_4-Xaa_5-Xaa_6-Xaa_7-Xaa_8-Xaa_9-Xaa_{10}-Xaa_{11}-Xaa_{12}-Xaa_{13}-Xaa_{14}-Xaa_{15}-Xaa_{16}-Xaa_{17}-Xaa_{18}-Xaa_{19}$$
 (IV)

wherein:

Xaa_a is proline; Xaa₁ is proline;

Xaa_b is glutamine or glutamic acid; Xaa₂ is arginine;

Xaa_c is threonine; Xaa₃ is cysteine;

Xaa_d is glycine; Xaa₄ is glycine;

Xaa_e is aspartic acid or glutamic acid; Xaa₅ is valine or asparagine;

Xaa_f is leucine; Xaa₆ is proline;

Xaag is aspartic acid; Xaa7 is aspartic acid;

Xaa_h is glutamine or serine; Xaa₈ is valine or leucine;

Xaa_i is asparagine or alanine; Xaa₉ is alanine or glycine;

Xaa_j is threonine; Xaa₁₀ is asparagine or arginine;

Xaa_k is isoleucine or leucine; Xaa₁₁ is tyrosine or phenylalanine;

Xaa_L is glutamic acid or lysine; Xaa₁₂ is asparagine or glutamine;

Xaa_m is threonine or alanine; Xaa₁₃ is phenylalanine or threonine;

Xaa_n is methionine; Xaa₁₄ is phenylalanine;

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Xaa_o is arginine;

Xaa₁₅ is proline or glutamic acid;

Xaa_p is lysine or threonine;

Xaa₁₆ is arginine or glycine;

Xaa₁₇ is lysine or aspartic acid;

Xaa₁₈ is proline or leucine; and

Xaa₁₉ is lysine.

- 22. (Original) The method of any one of claims 18-21, wherein the peptide comprises SEQ ID NO:1, SEQ ID NO:2, SEQ ID NO:3, SEQ ID NO:4, SEQ ID NO:5, SEQ ID NO:6, SEQ ID NO:7, SEQ ID NO:8, SEQ ID NO:9, SEQ ID NO:10, SEQ ID NO:11, SEQ ID NO:12, or SEQ ID NO:13.
- 23. (Original) The method of any one of claims 18-21, wherein the peptide comprises SEQ ID NO:11.
- 24. (Original) The method of any one of claims 18-21, wherein an apolar amino acid is methionine, glycine or proline.
- 25. (Original) The method of any one of claims 18-21, wherein a basic amino acid is histidine, lysine, arginine, 2,3-diaminopropionic acid, ornithine, homoarginine, ñaminophenylalanine, and 2,4-diaminobutyric acid.
- 26. (Original) The method of any one of claims 18-21, wherein a cysteine-like amino acid is cysteine, homocysteine, penicillamine, or â-methyl cysteine.
- 27. (Original) The method of any one of claims 18-21, wherein an aliphatic amino acid is alanine, valine, leucine, isoleucine, t-butylalanine, t-butylalanine, N-methylisoleucine, norleucine, N-methylvaline, cyclohexylalanine, â-alanine, N-methylglycine, or á-aminoisobutyric acid.

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28. (Original) The method of any one of claims 18-21, wherein an acidic amino acid is aspartic acid or glutamic acid.

- 29. (Original) The method of any one of claims 18-21, wherein a polar amino acid is asparagine, glutamine, serine, threonine, tyrosine, citrulline, N-acetyl lysine, methionine sulfoxide, or homoserine, or an apolar amino acid such as methionine, glycine or proline.
- 30. (Original) The method of any one of claims 18-21, wherein an aromatic amino acid is phenylalanine, tyrosine, tryptophan, phenylglycine, naphthylalanine, â-2-thienylalanine, 1,2,3,4-tetrahydro-isoquinoline-3-carboxylic acid, 4-chlorophenylalanine, 2-fluorophenylalanine, 3-fluorophenylalanine, 4-fluorophenylalanine, pyridylalanine, or 3-benzothienyl alanine.
- 31. (Original) The method of any one of claims 18-21, that further comprises locally administering the peptide to a tumor in a mammal.
- 32. (Original) The method of claim 31, wherein the tumor is metastatic, non-metastatic, vascularized, non-vascularized, hard or soft.
- 33. (Original) The method of any one of claims 18-21, wherein decreasing growth of chondrosarcoma cells decreases growth of conventional chondrosarcoma, myxoid chondrosarcoma, mesenchymal chondrosarcoma, clear cell chondrosarcoma, or dedifferentiated (spindle cell) chondrosarcoma.